

## Specific Cesium Transport via the *Escherichia coli* Kup (TrkD) K<sup>+</sup> Uptake System

DIRK BOSSEMEYER, ANDREAS SCHLÖSSER, AND EVERT P. BAKKER\*

Abteilung Mikrobiologie, Universität Osnabrück, Postfach 4469, D-4500 Osnabrück, Federal Republic of Germany

Received 3 November 1988/Accepted 24 December 1988

***Escherichia coli* cells which contain a functional Kup (formerly TrkD) system took up Cs<sup>+</sup> with a moderate rate and affinity. Kup is a separate K<sup>+</sup> uptake system with relatively little discrimination in the transport of the cations K<sup>+</sup>, Rb<sup>+</sup>, and Cs<sup>+</sup>. Regardless of the presence or absence of Kup, K<sup>+</sup>-replete cells took up Cs<sup>+</sup> primarily by a very low affinity mode, proportional to the ratio of the Cs<sup>+</sup> and K<sup>+</sup> concentrations in the medium.**

One of the striking consequences of the Chernobyl reactor catastrophe is the prolonged occurrence of radioactive cesium (Cs<sup>+</sup>) in animals, plants, and fungi. Higher animal cells show relatively little discrimination between the transport across the cytoplasmic membrane of K<sup>+</sup> (ionic radius, 0.133 nm) and the larger monovalent cations Rb<sup>+</sup> (radius, 0.148 nm) or Tl<sup>+</sup> (radius 0.143 nm) (5). The Na<sup>+</sup>/K<sup>+</sup>-ATPase catalyzes uptake of Cs<sup>+</sup> (radius, 0.169 nm) (24). By contrast, microorganisms and especially bacteria possess alkali cation transport systems that are much more selective for the different cations (13, 22). The constitutive K<sup>+</sup> uptake system Trk from *Escherichia coli* shows approximately a 10-fold discrimination between K<sup>+</sup> and Rb<sup>+</sup> or Tl<sup>+</sup> (1, 8, 20), and the inducible, high-affinity K<sup>+</sup> uptake ATPase Kdp is even more selective (8, 20). Since very little is known about Cs<sup>+</sup> accumulation by microorganisms besides what is in a few reports (7, 15) and since knowledge about such accumulation is essential for the estimation of the extent to which microorganisms may contribute to the prolonged retention of radioactive Cs<sup>+</sup> by soil, we determined which *E. coli* K<sup>+</sup> uptake systems accept Cs<sup>+</sup>. This is the only microorganism for which a number of well-defined K<sup>+</sup> transport mutants are available (10, 19, 23). We report here that Cs<sup>+</sup> is only taken up via the TrkD system. This observation supports the original but later abandoned view (cf. reference 19 with references 11 and 23) that the TrkD system is a separate K<sup>+</sup> uptake system. It possesses a low specificity for the alkali cations transported and is thereby clearly distinct from the two main K<sup>+</sup> uptake systems Trk and Kdp. To avoid future confusion with the Trk system we will refer to the TrkD system, which is the product of the *trkD* gene(s), as Kup (mnemonic for K<sup>+</sup> uptake).

**Bacterial strains and growth conditions.** All of the strains used are derivatives of *E. coli* K-12 kindly provided by W. Epstein, The University of Chicago, Chicago, Ill. The genotypes of these strains and the K<sup>+</sup> uptake systems expressed under the different growth conditions are listed in Table 1. The growth media were those described by Epstein and Kim (10).

**Plasmid pJG1.** The *trkD*-containing plasmid pJG1, described by Lopilato et al. (16), was obtained from W. Epstein. It consists of a 6.3-kilobase *EcoRI*-*HindIII* fragment cloned into the large *EcoRI*-*HindIII* fragment of pBR322 and contains the *trkD* gene (D. C. Dosch, Ph.D.

thesis, The University of Chicago, Chicago, Ill. 1985), as well as the *rbsA* gene (14).

**K<sup>+</sup> depletion and transport assays.** Growing cells were harvested by centrifugation after they had reached an optical density at 578 nm between 0.7 and 1.0. These cells were depleted of K<sup>+</sup> by Tris-EDTA treatment as described previously (4). Transport assays were carried out at 23 to 25°C with EDTA-treated cells suspended at 1 mg (dry wt) ml of medium<sup>-1</sup> consisting of 200 mM sodium HEPES (*N*-2-hydroxyethylpiperazine-*N'*-2-ethanesulfonic acid) (pH 7.5) and 10 mM glucose (4). After a preincubation of 15 min, uptake assays were started by the addition of KCl, <sup>86</sup>RbCl, or CsCl at the concentration indicated at the particular experiment. Cells from 1.0-ml samples were centrifuged through silicone oil. The K<sup>+</sup> and Cs<sup>+</sup> contents of the cell pellets were determined in a flame-photometer 700 (Eppendorf Geratebau, Hamburg, Federal Republic of Germany) and at 852.1 nm in the emission mode of an atomic absorption photometer 357 (Instrumentation Laboratory Inc., Lexington, Mass.), respectively. The <sup>86</sup>Rb<sup>+</sup> content of the cell pellets was determined in a 460C-liquid scintillation counter (Canberra-Packard, Frankfurt, Federal Republic of Germany).

**Cs<sup>+</sup> uptake by K<sup>+</sup> uptake mutants.** Figure 1 gives rates with which K<sup>+</sup>-depleted cells of the different K<sup>+</sup> transport mutants showed net Cs<sup>+</sup> uptake. Strains that possess a functional Kup system (i.e., FRAG-1, FRAG-5, or TK1001 [Fig. 1, closed symbols]) took up Cs<sup>+</sup> fairly rapidly ( $V_{\max}$ , 15 to 25  $\mu\text{mol min}^{-1}$  g [dry wt] of cells<sup>-1</sup>;  $K_m$ , 5 to 7 mM). Strains that only possess a Kdp system (strains TK2240 or TK2242) or Trk system (TK1001) took up Cs<sup>+</sup> as slowly as did strain TK2205, which is deleted for the Kdp system and carries point mutations in both the *trkA* and *trkD* genes (Fig. 1, open symbols).

**Enhanced Cs<sup>+</sup> uptake by cells carrying the *trkD* gene on a multicopy plasmid.** Cells that were either *kdp trkA trkD*, *kdp trkD trkE trkG*, or *kdp trkD trkG trkH* and contained the *trkD* gene on the multicopy plasmid pJG1 (16) took up K<sup>+</sup> several times faster than did cells that carried a functional *trkD* gene in the chromosome (D. Bossemeyer, I. R. Booth, and E. P. Bakker, Short Rep. Fifth Eur. Bioenergetics Conf. Aberystwyth, Wales, 1988, p. 199). The same was true for Cs<sup>+</sup> uptake (Fig. 1). Since Cs<sup>+</sup> is only taken up via the Kup system (see above), these results suggest that the *E. coli* chromosomal DNA of plasmid pJG1 alone is sufficient to encode a functional Kup system.

\* Corresponding author.

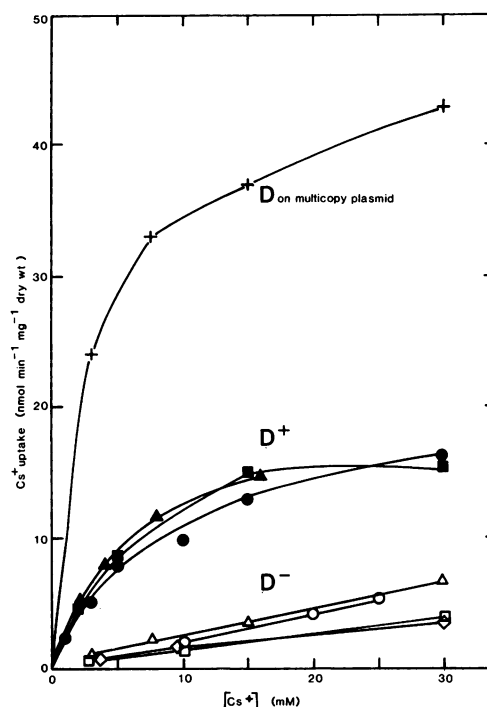
TABLE 1.  $K^+$  uptake mutants of *E. coli* K-12 and their properties<sup>a</sup>

Strain	Genotype <sup>b</sup>	[ $K^+$ ] during growth	$K^+$ transport systems expressed
FRAG-1	<i>gal</i>	30 mM	Trk, Kup
FRAG-5	<i>kdpABC5 gal</i>	5 mM	Trk, Kup
TK1001	<i>kdpABC5 trkD1 gal</i>	5 mM	Trk
TK1110	<i>kdpABC5 trkA405 gal</i>	5 mM	Kup
TK2205	<i>kdpABC5 trkD1 trkA405 gal</i>	115 mM	None <sup>c</sup>
TK2240	<i>trkA405 trkD1 nagA</i>	100 $\mu$ M <sup>d</sup>	Kdp
TK2242	<i>kdp-42 trkA405 trkD1 nagA</i>	5 mM	Kdp* <sup>e</sup>

<sup>a</sup> Genetic data are from references 10, 12, and 19.<sup>b</sup> All of the strains are also  $F^- lacZ rha thi$ .<sup>c</sup> The slow  $K^+$  uptake observed with this strain may be mediated by a fourth uptake system, TrkF (19).<sup>d</sup> Cells of strain TK2240 were grown overnight at 0.1 mM  $K^+$ . On the next day, they were diluted 1:20 into fresh medium without added  $K^+$ , but which was contaminated with about 25  $\mu$ M  $K^+$ . After growth had ceased, 25  $\mu$ M KCl was added to the suspension. After growth had ceased again,  $K^+$  was added at 100  $\mu$ M. The cells were harvested after growth had resumed.<sup>e</sup> The Kdp\* system possesses a greatly diminished affinity for  $K^+$  (12).

**Ion specificity of the Kup system.** That the Kup system accepts  $Cs^+$  suggests that it does not discriminate strongly between  $K^+$  and  $Cs^+$ . This differs from previous findings that the TrkD system distinguishes even more strongly between  $K^+$  and  $Rb^+$  than does the Trk system (20). We cannot confirm this earlier result. First, the kinetic parameters of  $K^+$  and  $Rb^+$  via Kup were almost identical (Table 2). Second, the uptake of tracer amounts of  $^{86}Rb^+$  by the cells in the presence of large amounts of either KCl or RbCl indicate that  $^{86}Rb$  uptake mimics bulk  $K^+$  uptake by the Kup system very well (results not shown). Table 2 also shows a large difference between the transport of  $K^+$  and  $Rb^+$  via the Kup system and that of  $Cs^+$  in the approximately 10-fold-higher  $K_m$  value of the latter. Thus, in contrast to Trk or Kdp, the Kup system does not strongly distinguish between the alkali cations  $K^+$ ,  $Rb^+$ , or even  $Cs^+$  (cf. Fig. 1 and Table 2 with references 1 and 20).

**Inhibition of  $K^+$  uptake by  $Cs^+$ .** Even a compound not translocated by a transport system may still inhibit the transport of the natural substrate. This effect was very small for  $Cs^+$  acting on  $K^+$  uptake by the Kdp system. A 30 mM concentration of  $Cs^+$  inhibited the net uptake of  $K^+$  (added at 2 mM) via the altered Kdp system (Table 1) of strain TK2242 by less than 10%. The effect of  $Cs^+$  on  $K^+$  uptake via the Trk system was larger with competitive inhibition and a  $K_i$  of 30 mM  $Cs^+$ . As expected, the strongest effect of  $Cs^+$  was observed on  $K^+$  uptake via the Kup system. The inhibition was competitive with a  $K_i$  of 7 mM  $Cs^+$  (results not shown).

FIG. 1. Cesium uptake by the Kup system. Cells of the *E. coli*  $K^+$  uptake mutants were tested for  $Cs^+$  uptake. Symbols: +, strain TK2205 containing the *trkD* gene on multicopy plasmid pJG1; ■, FRAG-1; ●, FRAG-5; ▲, TK1110; △, TK1001; ○, TK2205; □, TK2240; ◇, TK2242.

**$Cs^+$  uptake by  $K^+$ -loaded cells.** Both Kdp- and Trk-like bacterial  $K^+$  uptake systems shut off their activity when the cells become loaded with  $K^+$  (2, 17, 18). This effect is due to an increase in cell turgor during net  $K^+$  uptake (6, 9), which occurs because in this process  $K^+$  is partially exchanging against protons (3, 21). The decrease of Trk activity in  $K^+$ -loaded cells led to a remarkable effect when  $K^+$  and an excess of  $Cs^+$  were added simultaneously to  $K^+$ -depleted cells of either strain FRAG-5, which contains functional Trk and Kup systems (Fig. 2A), or TK1001, which only possesses a Trk system (Fig. 2B). In both situations, the cells were first rapidly loaded with  $K^+$ . However, subsequently the cells slowly exchanged  $K^+$  for  $Cs^+$ . From the exchange rate in strain TK1001, it can be calculated that in  $K^+$ -loaded cells the activity of the Trk system was as low as 3  $\mu$ mol  $min^{-1} g^{-1}$ , which is about 1 to 2% of the  $V_{max}$  for  $K^+$  uptake by  $K^+$ -depleted cells (1, 19). Hence, the effect of shutting off Trk activity during loading of the cells with  $K^+$  is much larger than has been recognized before (17, 18). However,

TABLE 2. Cation specificity of *E. coli*  $K^+$  uptake systems

System <sup>a</sup>	$K^+$ uptake <sup>b</sup>		$Rb^+$ uptake <sup>b</sup>		$Cs^+$ uptake <sup>b</sup>		References
	$K_m$	$V_{max}$	$K_m$	$V_{max}$	$K_m$	$V_{max}$	
Kdp	0.002	150	Low activity	Low activity	ND <sup>c</sup>	ND	19, 20, this work
Trk	0.9–1.5	190–500	0.35	18	ND	ND	1, 19, this work
Kup	$0.37 \pm 0.13^d$	$27 \pm 5^d$	$0.38 \pm 0.07^e$	$38 \pm 1^e$	$5 \pm 1^e$	$17 \pm 3^e$	This work

<sup>a</sup> Data from strains TK2240, TK1001, and TK1110 for the Kdp, Trk, and Kup systems, respectively.<sup>b</sup>  $K_m$  in millimolar;  $V_{max}$  in micromoles minute<sup>-1</sup> gram<sup>-1</sup> (dry wt) of cells.<sup>c</sup> ND, No activity detected.<sup>d</sup> Average from four experiments with standard deviations.<sup>e</sup> Average from two experiments with standard deviations.

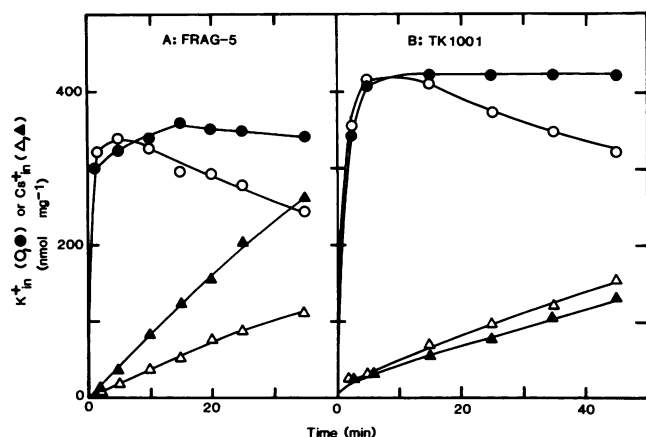


FIG. 2.  $K^+$ - $Cs^+$  exchange by *E. coli*. To glucose-metabolizing,  $K^+$ -depleted cells of strain FRAG-5 (A) or TK1001 (B) were added, at zero time, either 5 mM KCl (●), 30 mM CsCl (▲) or 5 mM KCl plus 30 mM CsCl (○, △); circles,  $K^+$  content; triangles,  $Cs^+$  content.

Mulder (M. M. Mulder, Ph.D. thesis, University of Amsterdam, Amsterdam, The Netherlands, 1988) drew a conclusion similar to ours.

**Very slow  $Cs^+$  uptake.** Cells that do not possess a functional Kup system took up  $Cs^+$  very slowly (Fig. 1). This  $Cs^+$  uptake nevertheless saturated at high  $Cs^+$  concentrations ( $V_{max}$ , 10 to 20  $\mu\text{mol min}^{-1} \text{g}^{-1}$ ;  $K_m$ , 100 to 150 mM; data not shown) and may either represent uptake via a fourth, very low affinity  $K^+$  uptake system, TrkF (19), or reflect a leak current through the cytoplasmic membrane driven by the internally negative membrane potential (19, 23). The second point illustrated by Fig. 2 is that in the presence of both  $K^+$  and  $Cs^+$ , the uptake of  $Cs^+$  by  $K^+$ -loaded cells occurred via this fourth system, since regardless of the presence or absence of a functional Kup system the two strains showed the same rate of  $Cs^+$  uptake (Fig. 2, open triangles). Since this system also shows almost no discrimination between  $K^+$  and  $Cs^+$  (data not shown), the extent of accumulation of radioactive  $Cs^+$  by wild-type,  $K^+$ -replete *E. coli* should be proportional to the ratio of the  $Cs^+$  and  $K^+$  concentrations present in the medium.

We thank Eva Limpinsel for expert technical assistance, G. Kooiker for help with the  $Cs^+$  determinations, and W. Epstein for strains and plasmid pJG1, as well as for helpful discussion.

This work was supported by grants from the Deutsche Forschungsgemeinschaft (SFB171, Teilprojekt C1) and from the Fonds der Chemischen Industrie.

#### LITERATURE CITED

1. Bakker, E. P. 1983. pH-dependent transport of rubidium by the constitutive potassium uptake system TrkA of *Escherichia coli* K-12. FEMS Microbiol. Lett. 16:229-233.
2. Bakker, E. P., and F. M. Harold. 1980. Energy coupling to

- potassium transport in *Streptococcus faecalis*. Interplay of ATP and the protonmotive force. J. Biol. Chem. 255:433-440.
3. Bakker, E. P., R. G. Kroll, and I. R. Booth. 1984. Potassium transport in *Escherichia coli*: sodium is not a substrate of the potassium uptake system TrkA. FEMS Microbiol. Lett. 23: 293-297.
4. Bakker, E. P., and W. E. Mangerich. 1981. Interconversion of components of the bacterial proton motive force by electrogenic potassium transport. J. Bacteriol. 147:820-826.
5. Bakker-Grunwald, T. 1979. Movement of thallous ion across the ascites cell membrane. J. Membr. Biol. 47:171-183.
6. Booth, I. R. 1985. Regulation of cytoplasmic pH in bacteria. Microbiol. Rev. 49:359-378.
7. Borst Pauwels, G. W. F. H. 1981. Ion transport in yeast. Biochim. Biophys. Acta 650:88-127.
8. Damper, P. D., W. Epstein, B. R. Rosen, and E. N. Sorensen. 1979. Thallous ion is accumulated by potassium transport systems in *Escherichia coli*. Biochemistry 18:4165-4169.
9. Epstein, W. 1986. Osmoregulation by potassium transport in *Escherichia coli*. FEMS Microbiol. Rev. 39:73-78.
10. Epstein, W., and B. S. Kim. 1971. Potassium transport loci in *Escherichia coli* K-12. J. Bacteriol. 108:639-644.
11. Epstein, W., and L. Laimins. 1980. Potassium transport in *Escherichia coli*: diverse systems with common control by osmotic forces. Trends Biochem. Sci. 5:21-23.
12. Epstein, W., V. Whitelaw, and J. Hesse. 1978. A  $K^+$  transport ATPase in *Escherichia coli*. J. Biol. Chem. 253:6666-6668.
13. Harold, F. M., and K. Altendorf. 1974. Cation transport in bacteria. Curr. Top. Membr. Transp. 5:1-50.
14. Iida, A., S. Harayama, T. Iino, and G. L. Hazelbauer. 1984. Molecular cloning and characterization of genes required for ribose transport and utilization in *Escherichia coli* K-12. J. Bacteriol. 158:674-682.
15. Jasper, P. 1978. Potassium transport system of *Rhodospseudomonas capsulata*. J. Bacteriol. 133:1314-1322.
16. Lopilato, J. E., J. L. Garwin, S. D. Emr, T. J. Silhavy, and J. R. Beckwith. 1984. D-Ribose metabolism in *Escherichia coli* K-12: genetics, regulation, and transport. J. Bacteriol. 158:665-673.
17. Meury, J., and A. Kepes. 1981. The regulation of potassium fluxes for the adjustment and maintenance of potassium levels in *Escherichia coli*. Eur. J. Biochem. 119:165-170.
18. Rhoads, D. B., and W. Epstein. 1978. Cation transport in *Escherichia coli*. IX. Regulation of K transport. J. Gen. Physiol. 72:283-295.
19. Rhoads, D. B., F. B. Waters, and W. Epstein. 1976. Cation transport in *Escherichia coli*. VIII. Potassium transport mutants. J. Gen. Physiol. 67:325-341.
20. Rhoads, D. B., A. Woo, and W. Epstein. 1977. Discrimination between  $Rb^+$  and  $K^+$  by *Escherichia coli*. Biochim. Biophys. Acta 469:45-51.
21. Schultz, S. G., W. Epstein, and A. K. Solomon. 1963. Cation transport in *Escherichia coli*. IV. Kinetics of net K uptake. J. Gen. Physiol. 47:329-346.
22. Silver, S. 1978. Transport of cations and anions, p. 221-324. In B. P. Rosen (ed.), Bacterial transport. Marcel Dekker, Inc., New York.
23. Walderhaug, M. O., D. C. Dosch, and W. Epstein. 1987. Potassium transport in bacteria, p. 84-130. In B. P. Rosen and S. Silver (ed.), Ion transport in procaryotes. Academic Press, Inc., New York.
24. Williams, R. J. P. 1970. Biochemistry of sodium, potassium, magnesium and calcium. Chem. Soc. Q. Rev. 24:331-365.